

RAW SEQUENCE LISTING

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Application Serial Number: 10/620,795A
Source: 1FW/b
Date Processed by STIC: 7/19/06

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DATE: 07/19/2006

PATENT APPLICATION: US/10/620,795A

TIME: 09:21:32

Input Set : F:\UTSK352.APP.txt

Output Set: N:\CRF4\07192006\J620795A.raw

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3 <110> APPLICANT: BARBOUR, ALAN G.
4     CARTER, CAROL
6 <120> TITLE OF INVENTION: A DIAGNOSTIC TEST FOR INFECTION WITH A SPIROCHETE BORNE
7     BY AMBLYOMMA AMERICANUM
9 <130> FILE REFERENCE: UTSK:352USC1
11 <140> CURRENT APPLICATION NUMBER: 10/620,795A
12 <141> CURRENT FILING DATE: 2003-07-14
14 <150> PRIOR APPLICATION NUMBER: 08/437,013
15 <151> PRIOR FILING DATE: 1995-05-08
17 <150> PRIOR APPLICATION NUMBER: 09/275,506
18 <151> PRIOR FILING DATE: 1999-03-24
20 <160> NUMBER OF SEQ ID NOS: 38
22 <170> SOFTWARE: PatentIn Ver. 2.1
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 641
26 <212> TYPE: DNA
27 <213> ORGANISM: Artificial Sequence
29 <220> FEATURE:
30 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
31     Primer
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35 attaacagag ttgctgatca ggctcaatac aaccagatgc atatgttatt taacaaatca 120
36 tctgctcaaa atgtaaaaac tgctgaagag cttggaatgc aacctgcaaa aattaatca 180
37 ccagcatcac taactggagc acaagcttca tggacattga gagttcaggt aggtgcaa 240
38 caggatgaag caattgctgt taatatcttc tcaactaatg ttgcaaactt ttttgggtga 300
39 gaagggtgttc aagcggctcc agctcaagag ggtgcacaac aggagggagt tcaaccagct 360
40 ccagctcaag gtgggattag ctctccaatt aatgttataa ctgctattga tgctaata 420
41 tcgcttataa agattgaaga tgctattaga atggtaactg atcaaagagc aaatcttggt 480
42 gctttccaaa atagacttga gtctgttaaa gctagcacag attatgctat tgaaaactta 540
43 aaagcgtctt atgctcaaat taaagatgca ataatgacag atgaaattgt agcatctaca 600
44 accaacagta ttttgacaca atctgcaatg gctatgattg c 641
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48 <211> LENGTH: 213
49 <212> TYPE: PRT
50 <213> ORGANISM: Artificial Sequence
52 <220> FEATURE:
53 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
54     Peptide
56 <400> SEQUENCE: 2
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60 Leu Thr Asp Glu Ile Asn Arg Val Ala Asp Gln Ala Gln Tyr Asn Gln

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61          20          25          30
63 Met His Met Leu Ser Asn Lys Ser Ser Ala Gln Asn Val Lys Thr Ala
64          35          40          45
66 Glu Glu Leu Gly Met Gln Pro Ala Lys Ile Asn Thr Pro Ala Ser Leu
67          50          55          60
69 Thr Gly Ala Gln Ala Ser Trp Thr Leu Arg Val Gln Val Gly Ala Asn
70 65          70          75          80
72 Gln Asp Glu Ala Ile Ala Val Asn Ile Phe Ser Thr Asn Val Ala Asn
73          85          90          95
75 Leu Phe Gly Gly Glu Gly Val Gln Ala Ala Pro Ala Gln Glu Gly Ala
76          100         105         110
78 Gln Gln Glu Gly Val Gln Pro Ala Pro Ala Gln Gly Gly Ile Ser Ser
79          115         120         125
81 Pro Ile Asn Val Thr Thr Ala Ile Asp Ala Asn Ala Ser Leu Thr Lys
82          130         135         140
84 Ile Glu Asp Ala Ile Arg Met Val Thr Asp Gln Arg Ala Asn Leu Gly
85 145         150         155         160
87 Ala Phe Gln Asn Arg Leu Glu Ser Val Lys Ala Ser Thr Asp Tyr Ala
88          165         170         175
90 Ile Glu Asn Leu Lys Ala Ser Tyr Ala Gln Ile Lys Asp Ala Ile Met
91          180         185         190
93 Thr Asp Glu Ile Val Ala Ser Thr Thr Asn Ser Ile Leu Thr Gln Ser
94          195         200         205
96 Ala Met Ala Met Ile
97          210
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101 <211> LENGTH: 1336
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103 <213> ORGANISM: Artificial Sequence
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106 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
107     Primer
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111 cgggtgagta acgcgtggat aatctgccta cgagatgggg ataactatta gaaataatag 120
112 ctaataccga ataaagtcaa ttgagttggt agttgatgaa aggaagcctt taaagcttcg 180
113 cttgtagatg agtctgcgtc ttattagcta gttggtaggg taagagccta ccaaggctat 240
114 gataagtaac cggcctgaga ggggtgatcg tcacactgga actgagatac ggtccagact 300
115 cctacgggag gcagcagcta agaattcttc gcaatgggag aaagcctgac ggagcgacac 360
116 tgcgtgaacg aagaaggctc aaagattgta aagttctttt ataaatgagg aataagcttt 420
117 gtaggaaatg acaagggtgat gacgttaatt tatgaataag ccccggttaa ttacgtgcca 480
118 gcagccgcgg taatacgtaa ggggcgagcg ttgttcggga tcattgggag taaaggggtga 540
119 gtaggcggat atgtaagtct atgtgtaaaa taccacggct caactgtgga actatgctag 600
120 aaactgcatg actagagtct gataggggaa gttagaattc ctggtgtaag ggtggaatct 660
121 gttgatatca ggaagaatac cagagggcga agcgaacctc taggtcaaga ctgacgctga 720
122 gtcacgaaag cgtagggagc aaacaggatt agataccctg gtagtctacg ctgtaaacga 780
123 tgcacacttg gtgttaatcg aaaggttagt accgaagcta acgtgttaag tgtgccgcct 840
124 ggggagtatg ctgcgaagag tgaaactcaa aggaattgac gggggccgcg acaagcgggtg 900
125 gagcatgtgg tttaattcga tgatacgcga ggaaccttac cagggttga catatacagg 960

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126 atatagttag agataactac tctccgtttg gggctctgtat acaggtgctg catggttgtc 1020
127 gtcagctcgt gctgtgaggt gttgggttaa gtcccgcaac gagcgcaacc cttgttgtct 1080
128 gttaccagca tgtaaagatg gggactcaga cgagactgcc ggtgataagc cggaggaagg 1140
129 tgaggatgac gtcaaatcat catggccctt atgtcctggg ctacacacgt gctacaatgg 1200
130 cctgtacaaa gcgatgcgaa acagtgatgt gaagcaaaac gcataaagca ggtctcagtc 1260
131 cagattgaag tctgaaactc gacttcatga agttggaatc gctagtaatc gtatatcaga 1320
132 atgatacggg gaatac 1336
135 <210> SEQ ID NO: 4
136 <211> LENGTH: 330
137 <212> TYPE: DNA
138 <213> ORGANISM: Artificial Sequence
140 <220> FEATURE:
141 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
142     Primer
144 <400> SEQUENCE: 4
145 aactgctgaa gagcttggaa tgcaacctgc aaaaattaat acaccagcat cactaactgg 60
146 agcacaagct tcatggacat tgagagttca ggtaggtgca aatcaggatg aagcaattgc 120
147 tgттаатatt ttctcaacta atgttgcaaa tctttttggg ggagaagggtg ttcaagcggc 180
148 tccagctcaa gaggggtgcac aacaggaggg agttcaacca gctccagctc aaggtgggat 240
149 tagctctcca attaatgtta caactgctat tgatgctaata gcacgctta caaagattga 300
150 agatgctatt agaatggtaa ctgatcaaag 330
153 <210> SEQ ID NO: 5
154 <211> LENGTH: 4
155 <212> TYPE: PRT
156 <213> ORGANISM: Artificial Sequence
158 <220> FEATURE:
159 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
160     Peptide
162 <400> SEQUENCE: 5
163 Gly Val Gln Ala
164     1
167 <210> SEQ ID NO: 6
168 <211> LENGTH: 9
169 <212> TYPE: DNA
170 <213> ORGANISM: Artificial Sequence
172 <220> FEATURE:
173 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
174     Primer
176 <400> SEQUENCE: 6
177 tctgctcaa 9
180 <210> SEQ ID NO: 7
181 <211> LENGTH: 12
182 <212> TYPE: DNA
183 <213> ORGANISM: Artificial Sequence
185 <220> FEATURE:
186 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
187     Primer
189 <400> SEQUENCE: 7
190 ggtgttcaag cg 12

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Input Set : F:\UTSK352.APP.txt

Output Set: N:\CRF4\07192006\J620795A.raw

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193 <210> SEQ ID NO: 8
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195 <212> TYPE: DNA
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198 <220> FEATURE:
199 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
200     Primer
202 <400> SEQUENCE: 8
203 gttcaaccag ct                                     12
206 <210> SEQ ID NO: 9
207 <211> LENGTH: 22
208 <212> TYPE: DNA
209 <213> ORGANISM: Artificial Sequence
211 <220> FEATURE:
212 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
213     Primer
215 <400> SEQUENCE: 9
216 aacagctgaa gagcttggaa tg                         22
219 <210> SEQ ID NO: 10
220 <211> LENGTH: 26
221 <212> TYPE: DNA
222 <213> ORGANISM: Artificial Sequence
224 <220> FEATURE:
225 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
226     Primer
228 <400> SEQUENCE: 10
229 cgataatctt actattcact agtttc                       26
232 <210> SEQ ID NO: 11
233 <211> LENGTH: 24
234 <212> TYPE: DNA
235 <213> ORGANISM: Artificial Sequence
237 <220> FEATURE:
238 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
239     Primer
241 <400> SEQUENCE: 11
242 acatattcag atgcagacag aggt                         24
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246 <211> LENGTH: 24
247 <212> TYPE: DNA
248 <213> ORGANISM: Artificial Sequence
250 <220> FEATURE:
251 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
252     Primer
254 <400> SEQUENCE: 12
255 tgttagacgt taccgttact aacg                         24
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259 <211> LENGTH: 20
260 <212> TYPE: DNA
261 <213> ORGANISM: Artificial Sequence

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263 <220> FEATURE:
264 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
265     Primer
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273 <212> TYPE: DNA
274 <213> ORGANISM: Artificial Sequence
276 <220> FEATURE:
277 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
278     Primer
280 <400> SEQUENCE: 14
281 catatagtct tactatgccca cttag                            25
284 <210> SEQ ID NO: 15
285 <211> LENGTH: 31
286 <212> TYPE: PRT
287 <213> ORGANISM: Artificial Sequence
289 <220> FEATURE:
290 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
291     Peptide
293 <400> SEQUENCE: 15
294 Leu Arg Val Gln Val Gly Ala Asn Gln Asp Glu Ala Ile Ala Val Asn
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297 Ile Phe Ser Thr Asn Val Ala Asn Leu Phe Gly Gly Glu Gly Val
298       20           25           30
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302 <211> LENGTH: 16
303 <212> TYPE: PRT
304 <213> ORGANISM: Artificial Sequence
306 <220> FEATURE:
307 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
308     Peptide
310 <400> SEQUENCE: 16
311 Gln Ala Ala Pro Ala Gln Glu Gly Ala Gln Gln Glu Gly Val Gln Pro
312   1           5           10           15
315 <210> SEQ ID NO: 17
316 <211> LENGTH: 20
317 <212> TYPE: PRT
318 <213> ORGANISM: Artificial Sequence
320 <220> FEATURE:
321 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
322     Peptide
324 <400> SEQUENCE: 17
325 Ala Pro Ala Gln Gly Gly Ile Ser Ser Pro Ile Asn Val Thr Thr Ala
326   1           5           10           15
328 Ile Asp Ala Asn
329       20
332 <210> SEQ ID NO: 18

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VERIFICATION SUMMARY

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